Application No. 10/795,795
Response and Amendment dated December 27, 2005
Reply to Office action of September 27, 2005
Docket Number 27433/04016

## AMENDMENTS TO THE CLAIMS

## Listing of the Claims:

27. (currently amended) A method for treating human subjects with blood clotting disorders, comprising:

administering a pharmaceutical composition to the human subjects,

wherein the pharmaceutical composition comprises a peptide <u>having a total length</u> of from 3 to 10 amino acids in total length,

wherein said peptide comprises a sequence of between 3 and 10 amino acids, which is identical to a sequence of from 3 to 10 consecutive amino acids found within amino acids 32216 to 33125 or 35246 to 35650 of the human blood clotting factor Va (SEQ ID NO.0: 1), wherein SEQ ID No: 1 represents amino acids 307 to 356 of the human blood clotting factor Va and

wherein the peptide exhibits an IC<sub>50</sub> of between 50 nM to 500  $\mu$ M for inhibition of prothrombinase.

- 28. (currently amended) The method of claim 27 wherein the peptide comprises at least 5 amino acids which are identical to a sequence of consecutive amino acids found within amino acids 32216 to 33125 or 35246 to 35650 of the human blood clotting factor Va (SEQ ID NO:0:1), wherein SEQ ID No: 1 represents amino acids 307 to 356 of the human blood clotting factor Va.
- 29. (currently amended) The method of claim 27 wherein the peptide comprises at least 7 amino acids which are identical to a sequence of consecutive amino acids found within amino acids 32216 to 33125 of the human blood clotting factor Va (SEQ ID NO.0: 1), wherein SEQ ID No: 1 represents amino acids 307 to 356 of the human blood clotting factor Va.
- 30. (currently amended) The method of claim 27 wherein the peptide comprises at least 10 amino acids which are identical to a sequence of consecutive amino acids found within amino acids 32216 to 33125 of the human blood clotting factor-Va-(SEQ ID NO.0: 1), wherein SEQ ID NO: 1 represents amino acids 307 to 356 of the human blood clotting factor Va.

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- 31. (currently amended) The method of claim 27, wherein the sequence of then peptide is selected from the group consisting of SEQ ID NO.0: 6 and SEQ ID NO.0: 12.
- 32. (previously presented) The method of claim 27, wherein at least one amino acid within said peptide is a non-naturally occurring amino acid.
- 33. (previously presented) The method of claim 27, wherein at least one amino acid within said peptide is a D-amino acid.
- 34. (currently amended) The method of claim 271, wherein at least two amino acids in said sequence are joined by non-hydrolyzable peptide bonds.
- 35. (currently amended) The peptide of claim 427, wherein said peptide is a cyclized peptide.